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<p>(54) Title: FLK-1 BINDING PROTEIN</p> <p>(57) Abstract</p> <p>A protein designated flk-1bp binds the vascular endothelial cell surface receptor flk-1. Isolated DNA encoding flk-1bp is provided, along with expression vectors and transformed host cells useful in producing flk-1bp polypeptides. Antibodies that are immunoreactive with flk-1bp are generated using the polypeptides disclosed herein.</p>		

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TITLE**flk-1 BINDING PROTEIN****BACKGROUND OF THE INVENTION**

10 The receptor tyrosine kinase (RTK) family includes receptors that are crucial for the growth and differentiation of a variety of cell types (Yarden and Ullrich, *Ann. Rev. Biochem.* 57:433-478, 1988; Ullrich and Schlessinger, *Cell* 61:243-254, 1990). Several receptor tyrosine kinases, and growth factors that bind thereto, have been suggested to play a role in angiogenesis, although some may promote angiogenesis indirectly (Mustonen and
15 Alitalo, *J. Cell Biol.* 129:895-898, 1995).

 One such receptor tyrosine kinase, known as fetal liver kinase 1 (flk-1), is a member of the type III subclass of RTKs. An alternative designation for human flk-1 is kinase insert domain-containing receptor (KDR) (Terman et al., *Oncogene* 6:1677-83, 1991), and the rat homolog has been termed TKr-C (Sarzani et al., *Biochem. Biophys. Res. Comm.* 186:706-714, 1992). DNAs encoding mouse, rat and human flk-1 have been
20 isolated, and the nucleotide and encoded amino acid sequences reported (Matthews et al., *Proc. Natl. Acad. Sci. USA*, 88:9026-30, 1991; Terman et al., 1991, *supra*; Terman et al., *Biochem. Biophys. Res. Comm.* 187:1579-86, 1992; Sarzani et al., *supra*; and Millauer et al., *Cell* 72:835-846, 1993).

25 The type III subclass RTK designated *fms*-like tyrosine kinase-1 (flt-1) is related to flk-1 (DeVries et al., *Science* 255:989-991, 1992; Shibuya et al., *Oncogene* 5:519-524, 1990). Flt-1 is believed to be essential for endothelial organization during vascular development. Flt-1 expression is associated with early vascular development in mouse embryos, and with neovascularization during wound healing (Mustonen and Alitalo,
30 *supra*). Expression of flt-1 in adult organs suggests an additional function for this receptor that is not related to cell growth (Mustonen and Alitalo, *supra*).

 Another RTK that is related to flt1 and flk1 is flt4 (Galland et al., *Oncogene* 8:1233-40, 1993; Pajusola et al., *Oncogene* 8:2931-37, 1993). Features shared by these three receptors include the seven immunoglobulin-like domains in their extracellular region.
35 The amino acid sequence of flt4 exhibits significant homology with the sequences of flt1 and flk1, especially in the tyrosine kinase domain (Galland et al., *supra*). Unlike flt-1 and flk-1, however, a precursor form of flt-4 is cleaved during post-translational processing to

form two disulfide-linked polypeptides (Pajusola et al., *supra*). Studies of Flt-4 expression during development support the theory of venous origin of lymphatic vessels (Kaipainen et al., *Proc. Natl. Acad. Sci. USA* 92:3566-70, April, 1995).

Given the crucial role of endothelial cells in angiogenesis, growth factors that act on
5 endothelial cells are of particular interest for studies of the regulation of vascularization. One such factor is vascular endothelial cell growth factor (VEGF), which binds to both flk-1 and flt-1 with relatively high affinity and is mitogenic toward vascular endothelial cells (Terman et al., 1992, *supra*; Mustonen et al. *supra*; DeVries et al., *supra*). VEGF does not
10 bind to flt4 (Pajusola et al., *supra*). The studies reported in Millauer et al., *supra*, suggest that VEGF and flk-1 are a ligand-receptor pair that play an important role in the formation and sprouting of blood vessels, termed vasculogenesis and angiogenesis, respectively.

Different forms of VEGF arising from alternative splicing of mRNA have been reported, including the four species described by Ferrara et al. (*J. Cell. Biochem.* 47:211-218, 1991). Both secreted and predominantly cell-associated species of VEGF were
15 identified by Ferrara et al. *supra*, and the protein is known to exist in the form of disulfide linked dimers.

Placenta growth factor (PlGF) has an amino acid sequence that exhibits significant homology to the VEGF sequence (Park et al., *J. Biol. Chem.* 269:25646-54, 1994; Maglione et al. *Oncogene* 8:925-31, 1993). As with VEGF, different species of PlGF
20 arise from alternative splicing of mRNA, and the protein exists in dimeric form (Park et al., *supra*). PlGF binds flt-1 with high affinity, but not flk-1 (Park et al., *supra*). PlGF potentiates the mitogenic effect of VEGF on endothelial cells when VEGF is present at low concentrations, but has no detectable effect when VEGF is present at higher concentrations (Park et al., *supra*).

25 Studies of growth factors and receptors that are believed to regulate angiogenesis include those discussed above. Investigation into the existence and identity of other such receptors, and proteins that bind thereto, is desirable. Identifying such proteins would provide additional means for elucidating the effects of various ligand-receptor signaling systems on development and differentiation of the vascular system, as well as providing
30 further insight into, and means for, regulation of such biological processes.

Inhibiting angiogenesis is desirable in certain clinical situations (e.g., to suppress growth and metastasis of solid tumors, or in treating rheumatoid arthritis), whereas promoting vascularization is beneficial for treating other conditions (e.g., wound healing). Consequently, molecules that promote angiogenesis by transducing signals through the
35 above-discussed receptors, and molecules capable of inhibiting such signal transduction, are both of interest.

SUMMARY OF THE INVENTION

The present invention provides a novel protein designated flk-1 binding protein (flk-1bp), as well as isolated DNA encoding the novel protein and expression vectors comprising the isolated DNA. A method for producing flk-1bp involves culturing host
5 cells transformed with the recombinant expression vectors under conditions appropriate for expression of flk-1bp, then recovering the protein from the culture. Secreted flk-1bp polypeptides are recovered from the culture medium.

The flk-1bp protein is able to bind to a cell surface receptor known as human flk-1 or KDR. This receptor is expressed on cell types that include vascular endothelial cells.
10 Antibodies that specifically bind flk-1bp polypeptides are also provided.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 presents an alignment of the amino acid sequences of three proteins, which are a human flk-1 binding protein of the present invention (flk-1bp) (SEQ ID NO:2), a
15 human vascular endothelial cell growth factor (VEGF) (SEQ ID NO:4), and a human placenta growth factor (PLGF) (SEQ ID NO:5). Gaps were introduced as necessary to maximize the alignment. Residues found in at least two of the sequences are boxed.

DETAILED DESCRIPTION OF THE INVENTION

20 A novel protein designated Flk-1bp is provided herein, along with DNA encoding flk-1bp and recombinant expression vectors comprising the flk-1bp DNA. A method for producing recombinant flk-1bp polypeptides involves cultivating host cells transformed with the expression vectors under conditions appropriate for expression of flk-1bp, and recovering the expressed flk-1bp.

25 The present invention also provides antibodies that specifically bind flk-1bp proteins. In one embodiment, the antibodies are monoclonal antibodies.

Isolation of a DNA encoding a human flk-1bp is described in example 1 below. The nucleotide sequence of the human flk-1bp DNA isolated in example 1 is presented in SEQ ID NO:1, and the amino acid sequence encoded thereby is presented in SEQ ID NO:2.
30 This human flk-1bp is a secreted protein that comprises an N-terminal signal peptide (amino acids -20 through -1 of SEQ ID NO:2) when initially synthesized. The signal peptide is cleaved during secretion of the protein from the cell, yielding a mature protein comprising amino acids 1 to 399 of SEQ ID NO:2.

E. coli strain DH10B cells transformed with a recombinant vector containing this
35 human flk-1bp DNA were deposited with the American Type Culture Collection on September 1, 1995, and assigned accession no. ATCC 69897. The deposit was made under the terms of the Budapest Treaty. As described in more detail in example 1, the

recombinant vector in the deposited strain contains human flk-1bp DNA that includes the entire coding region shown in SEQ ID NO:1.

In one embodiment, the present invention provides a flk-1bp protein encoded by the flk-1bp DNA insert of the vector in deposited strain ATCC 69897. flk-1bp protein
5 expressed by a host cell transformed with an expression vector containing the flk-1bp DNA insert found in the vector in strain ATCC 69897 is provided.

The flk-1bp of SEQ ID NO:2 contains an N-terminal signal peptide, and is capable of being secreted from suitable host cells in which it is expressed. The flk-1bp thus may be recovered from the culture medium, which facilitates purification.

10 Vascular endothelial cell growth factor (VEGF) binds to the receptor tyrosine kinase flk-1 (DeVries et al. *supra*). As demonstrated in example 3, human flk-1bp demonstrated the ability to compete with VEGF for binding to human flk-1.

Figure 1 presents an alignment of the amino acid sequences of three proteins, which are a human flk-1 binding protein of the present invention (flk-1bp) (SEQ ID NO:2), a
15 human vascular endothelial cell growth factor (VEGF) (SEQ ID NO:4), and a human placenta growth factor (PLGF) (SEQ ID NO:5). Gaps were introduced as necessary to maximize the alignment. Residues found in at least two of the sequences are boxed.

As described in example 1, a search of a sequence databank was performed, using a human VEGF amino acid sequence as the query sequence. An expressed sequence tag
20 (EST) file, GenBank accession number Z44272, was identified by the search. The 299-nucleotide sequence presented in file no. Z44272 corresponds to nucleotides 660 to 958 of SEQ ID NO:1, with the exception of nucleotide number 733, which was not identified in file no. Z44272. The NCBI/GenBank record does not disclose any polypeptide encoded by the EST of file no. Z44272, and does not indicate what the reading frame, if any, might
25 be.

One embodiment of the present invention is directed to mature human flk-1bp protein characterized by the N-terminal amino acid sequence Gly-Pro-Arg-Glu-Ala-Pro-Ala-Ala-Ala-Ala-Ala-Phe-Glu-Ser-Gly- (amino acids 1-15 of SEQ ID NO:2). As
30 described in example 1, the present invention provides mature flk-1bp protein characterized by a calculated molecular weight of 44,823 daltons and an isoelectric point of 8.678.

The flk-1bp polypeptides of the present invention include polypeptides having amino acid sequences that differ from, but are highly homologous to, that presented in SEQ ID NO:2. Examples include, but are not limited to, homologs derived from other
35 mammalian species (which can be identified by cross-species hybridization using human flk-1bp DNA as a probe), variants (both naturally occurring variants and those generated by recombinant DNA technology), and flk-1bp fragments that retain a desired biological activity. Such polypeptides exhibit a biological activity of the flk-1bp proteins of SEQ ID

NO:2, and preferably comprise an amino acid sequence that is at least 80% identical (most preferably at least 90% identical) to the amino acid sequence presented in SEQ ID NO:2. These embodiments of the present invention are described in more detail below.

flk-1bp fragments may be prepared by any of a number of conventional techniques.

- 5 A DNA sequence encoding a desired flk-1bp fragment may be subcloned into an expression vector for production of the flk-1bp fragment. The desired flk-1bp-encoding DNA fragment may be chemically synthesized using known techniques. DNA fragments also may be produced by restriction endonuclease digestion of a full length cloned DNA sequence, and isolated by electrophoresis on agarose gels. If necessary, oligonucleotides that reconstruct the 5' or 3' terminus to a desired point may be ligated to a DNA fragment
10 generated by restriction enzyme digestion. Such oligonucleotides may additionally contain a restriction endonuclease cleavage site upstream of the desired coding sequence, and position an initiation codon (ATG) at the N-terminus of the coding sequence.

- The well known polymerase chain reaction (PCR) procedure also may be employed
15 to isolate and amplify a DNA sequence encoding a desired protein fragment. Oligonucleotides that define the desired termini of the DNA fragment are employed as 5' and 3' primers. The oligonucleotides may additionally contain recognition sites for restriction endonucleases, to facilitate insertion of the amplified DNA fragment into an expression vector. PCR techniques are described in Saiki et al., *Science* 239:487 (1988);
20 *Recombinant DNA Methodology*, Wu et al., eds., Academic Press, Inc., San Diego (1989), pp. 189-196; and *PCR Protocols: A Guide to Methods and Applications*, Innis et al., eds., Academic Press, Inc. (1990).

- The flk-1bp signal peptide discussed above is identified in accordance with conventional criteria for identifying that type of hydrophobic domain. Computer programs
25 that predict the site of cleavage of a signal peptide are available. However, more than one possible cleavage site may be identified by such computer programs, and it is recognized in the pertinent field that cleavage can occur at sites other than those predicted. Signal peptide cleavage sites that are less likely, but possible, for flk-1bp occur after residue -6 (in which case the mature protein contains amino acids -5 to 399 of SEQ ID NO:2) or after residue -5
30 (in which case the mature protein contains amino acids -4 to 399 of SEQ ID NO:2). As will be understood by the skilled artisan, a protein preparation can comprise a mixture of protein molecules having different N-terminal amino acids, due to cleavage of the signal peptide at more than one site.

- In addition, post-translational processing can vary according to the particular
35 expression system employed. Consequently, the N- or C-terminal amino acid of a recombinant protein may vary according to the type of host cells in which the protein was expressed, for example. Particular embodiments include but are not limited to flk-1bp

proteins having an N-terminal amino acid selected from any of amino acids -5, -4, -3, -2, -1, 1, 2, 3, 4, and 5; and a C-terminal amino acid selected from amino acids 395 to 399 of SEQ ID NO:2.

5 Mature flk-1bp provided herein includes such naturally occurring variants, in which the N-terminal amino acid is other than the residue at position 1, or the C-terminal amino acid is other than amino acid 399, of SEQ ID NO:2. Other naturally occurring flk-1bp variants encompassed by the present invention are those arising from alternative mRNA splicing events (since flk-1bp is encoded by a multi-exon gene). Other naturally occurring variants are allelic variants.

10 The flk-1bp DNA of the present invention includes cDNA, chemically synthesized DNA, DNA isolated by PCR, genomic DNA, and combinations thereof. Genomic flk-1bp DNA may be isolated by hybridization to the flk-1bp cDNA disclosed herein using standard techniques. RNA transcribed from the flk-1bp DNA is also encompassed by the present invention.

15 Certain embodiments of the present invention provide isolated DNA comprising a nucleotide sequence selected from the group consisting of nucleotides 188 to 1444 of SEQ ID NO:1 (human flk-1bp coding region) and nucleotides 248 to 1444 of SEQ ID NO:1 (encoding a mature human flk-1bp protein). DNAs encoding biologically active fragments of the protein of SEQ ID NO:2 are also provided.

20 Due to degeneracy of the genetic code, two DNA sequences may differ, yet encode the same amino acid sequence. The present invention thus provides isolated DNA sequences encoding biologically active flk-1bp, selected from DNA comprising the coding region of a native flk-1bp cDNA, or fragments thereof, and DNA which is degenerate as a result of the genetic code to a native flk-1bp-encoding DNA sequence.

25 Also provided herein are purified flk-1bp polypeptides, both recombinant and non-recombinant. Variants and derivatives of native flk-1bp proteins that retain a desired biological activity are also within the scope of the present invention. In one embodiment, the biological activity of a flk-1bp variant is essentially equivalent to a desired biological activity of a native flk-1bp protein. One biological activity of human flk-1bp is the ability to bind human flk-1. The ability of a flk-1bp polypeptide (including variants thereof) to
30 bind flk-1 can be determined in a conventional binding assay.

It is possible that flk-1bp can bind to other receptors that are related to flk-1, such as flt-1. The ability of flk-1bp to bind to flt-1 can be determined in binding assays. As an alternative to binding assays, assays for a biological effect resulting from interaction of flk-1bp with a given receptor may be employed. Examples include signal transduction assays.
35 Phosphorylation of a receptor upon binding flk-1bp may be detected by known procedures, which may include use of commercially available anti-phosphotyrosine antibodies

(Chemicon International, Temecula, CA). Additional receptors with which flk-1bp interacts thus may be identified.

A receptor tyrosine kinase designated flt-4 has been described (Galland et al., *Oncogene* 8:1233-40, 1993; Pajusola et al., *Oncogene* 8:2931-37, 1993; Kaipainen et al.,
5 *Proc. Natl. Acad. Sci. USA* 92:3566-70, April, 1995). The flk-1bp of the present invention binds, and induces phosphorylation of, flt-4.

flk-1bp variants may be obtained by mutations of native flk-1bp nucleotide sequences, for example. An flk-1bp variant, as referred to herein, is a polypeptide substantially homologous to a native flk-1bp, but which has an amino acid sequence
10 different from that of native flk-1bp because of one or a plurality of deletions, insertions or substitutions. flk-1bp-encoding DNA sequences of the present invention encompass sequences that comprise one or more additions, deletions, or substitutions of nucleotides when compared to a native flk-1bp DNA sequence, but that encode an flk-1bp protein that is essentially biologically equivalent to a native flk-1bp protein.

15 The variant amino acid or DNA sequence preferably is at least 80% identical to a native flk-1bp sequence, most preferably at least 90% identical. In particular embodiments, DNA or amino acid sequences are at least 95%, 96%, 97%, 98%, or 99% identical to the sequence presented in SEQ ID NOS:1 or 2. The degree of homology (percent identity) between a native and a mutant sequence may be determined, for example, by comparing the
20 two sequences using computer programs commonly employed for this purpose. One suitable program is the GAP computer program, version 6.0, described by Devereux et al. (*Nucl. Acids Res.* 12:387, 1984) and available from the University of Wisconsin Genetics Computer Group (UWGCG). The GAP program utilizes the alignment method of Needleman and Wunsch (*J. Mol. Biol.* 48:443, 1970), as revised by Smith and Waterman
25 (*Adv. Appl. Math* 2:482, 1981). Briefly, the GAP program defines identity as the number of aligned symbols (i.e., nucleotides or amino acids) which are identical, divided by the total number of symbols in the shorter of the two sequences. The preferred default parameters for the GAP program include: (1) a unary comparison matrix (containing a value of 1 for identities and 0 for non-identities) for nucleotides, and the weighted
30 comparison matrix of Gribskov and Burgess, *Nucl. Acids Res.* 14:6745, 1986, as described by Schwartz and Dayhoff, eds., *Atlas of Protein Sequence and Structure*, National Biomedical Research Foundation, pp. 353-358, 1979; (2) a penalty of 3.0 for each gap and an additional 0.10 penalty for each symbol in each gap; and (3) no penalty for end gaps.

35 DNA sequences encoding flk-1bp (including fragments and variants thereof) include DNA sequences that will hybridize to a native flk-1bp DNA under conditions of moderate or severe stringency, wherein the encoded flk-1bp retains a desired biological

activity (e.g., the ability to bind flk-1). Moderate stringency hybridization conditions refer to conditions described in, for example, Sambrook et al. *Molecular Cloning: A Laboratory Manual*, 2 ed. Vol. 1, pp. 1.101-104, Cold Spring Harbor Laboratory Press (1989).

5 Conditions of moderate stringency, as defined by Sambrook et al., include use of a prewashing solution of 5 X SSC, 0.5% SDS, 1.0 mM EDTA (pH 8.0) and hybridization conditions of about 55°C, 5 X SSC, overnight. Conditions of severe stringency include higher temperatures of hybridization and washing, as is understood by one skilled in the field of molecular biology. In one embodiment of the invention, a flk-1bp is encoded by a DNA sequence that will hybridize under severely stringent conditions to a DNA sequence
10 comprising nucleotides 188 to 1444 of SEQ ID NO:1 (the coding region of the flk-1bp DNA of SEQ ID NO:1). The severely stringent conditions include hybridization at 68°C followed by washing in 0.1X SSC/0.1% SDS at 63-68°C. flk-1bp polypeptides encoded by such hybridizing DNA sequences are provided herein.

Alterations of the native amino acid sequence may be accomplished by any of a
15 number of known techniques. Mutations can be introduced at particular loci by synthesizing oligonucleotides containing a mutant sequence, flanked by restriction sites enabling ligation to fragments of the native sequence. Following ligation, the resulting reconstructed sequence encodes an analog having the desired amino acid insertion, substitution, or deletion.

20 Alternatively, oligonucleotide-directed site-specific mutagenesis procedures can be employed to provide an altered gene having particular codons altered according to the substitution, deletion, or insertion required. Techniques for making such alterations include those disclosed by Walder et al. (*Gene* 42:133, 1986); Bauer et al. (*Gene* 37:73, 1985); Craik (*BioTechniques*, January 1985, 12-19); Smith et al. (*Genetic Engineering: Principles and Methods*, Plenum Press, 1981); and U.S. Patent Nos. 4,518,584 and
25 4,737,462, which are incorporated by reference herein.

Variants may comprise conservatively substituted sequences, meaning that one or more amino acid residues of a native flk-1bp polypeptide are replaced by different residues, but that the conservatively substituted flk-1bp polypeptide retains a desired biological
30 activity that is essentially equivalent to that of a native flk-1bp polypeptide. Examples of conservative substitutions include substitution of amino acids that do not alter the secondary and/or tertiary structure of flk-1bp. Other examples involve substitution of amino acids outside of the receptor-binding domain, when the desired biological activity is the ability to bind to a particular receptor.

35 A given amino acid may be replaced by a residue having similar physiochemical characteristics, e.g., substituting one aliphatic residue for another (such as Ile, Val, Leu, or Ala for one another), or substitution of one polar residue for another (such as between Lys

and Arg; Glu and Asp; or Gln and Asn). Other such conservative substitutions, e.g., substitutions of entire regions having similar hydrophobicity characteristics, are well known. flk-1bp polypeptides comprising conservative amino acid substitutions may be tested in one of the assays described herein to confirm that a desired biological activity of a native flk-1bp is retained. DNA sequences encoding flk-1bp polypeptides that contain such conservative amino acid substitutions are encompassed by the present invention.

Residues in flk-1bp that are conserved, in that identical residues in VEGF or PlGF (or both) are aligned with these flk-1bp residues in Figure 1, are amino acids -20, -7, 111, 115, 119, 123, 132, 134, 136, 137, 141, 142, 144-146, 149, 151, 153, 168, 182, 189, 207, 323, 335, 338, 340, 342, 360, 361, and 363 of SEQ ID NO:2. Advantageously, these conserved amino acids are not altered when generating conservatively substituted sequences or other variants of a native flk-1bp.

The conserved amino acids in mature flk-1bp are clustered into two regions, namely, from amino acids 111 to 207, and from amino acids 323 to 363 of SEQ ID NO:2. flk-1bp fragments that contain the conserved amino acids are encompassed by the present invention. Such fragments may contain amino acids 111 to 207 of SEQ ID NO:2, or amino acids 323 to 363 of SEQ ID NO:2, or both.

Figure 1 shows that there are no sequences in VEGF or PlGF that are aligned with amino acids 27 through 105, amino acids 209 through 322, and amino acids 369 through 399 of SEQ ID NO:2. Fragments of flk-1bp that lack from one to all of residues 27 through 105, 209 through 322, or 369 through 399 of SEQ ID NO:2 are encompassed by the present invention.

VEGF binds flk-1 and flt-1, whereas PlGF binds flt-1 but not flk-1. It is noteworthy that PlGF contains the conserved amino acids that are clustered in the region aligned with amino acids 111 to 207 of flk-1bp, but lacks the conserved amino acids found between amino acids 323 to 363 of flk-1bp. VEGF contains both groups of conserved amino acids.

flk-1bp may be modified to create flk-1bp derivatives by forming covalent or aggregative conjugates with other chemical moieties, such as glycosyl groups, lipids, phosphate, acetyl groups and the like. Covalent derivatives of flk-1bp may be prepared by linking the chemical moieties to functional groups on flk-1bp amino acid side chains or at the N-terminus or C-terminus of a flk-1bp polypeptide or the extracellular domain thereof.

Other derivatives of flk-1bp within the scope of this invention include covalent or aggregative conjugates of flk-1bp or its fragments with other proteins or polypeptides, such as by synthesis in recombinant culture as N-terminal or C-terminal fusions. For example, the conjugate may comprise a heterologous signal or leader polypeptide sequence in place of the native signal peptide at the N-terminus of a flk-1bp polypeptide. The signal or

leader peptide co-translationally or post-translationally directs transfer of the conjugate from its site of synthesis to a site inside or outside of the cell membrane or cell wall. Another example is a fusion protein comprising flk-1bp and a polypeptide that promotes oligomerization. Examples of such fusion proteins are described in more detail below.

5 flk-1bp polypeptide fusions can comprise peptides added to facilitate purification and identification of flk-1bp. Such peptides include, for example, poly-His or the antigenic identification peptides described in U.S. Patent No. 5,011,912 and in Hopp et al., *BioTechnology* 6:1204, 1988. One such peptide is the FLAG® peptide, Asp-Tyr-Lys-Asp-Asp-Asp-Lys (DYKDDDDK) (SEQ ID NO:3), which is highly antigenic and
10 provides an epitope reversibly bound by a specific monoclonal antibody, thus enabling rapid assay and facile purification of expressed recombinant protein. This sequence is also specifically cleaved by bovine mucosal enterokinase at the residue immediately following the Asp-Lys pairing. Fusion proteins capped with this peptide may also be resistant to intracellular degradation in *E. coli*. A murine hybridoma designated 4E11 produces a
15 monoclonal antibody that binds the peptide DYKDDDDK (SEQ ID NO:3) in the presence of certain divalent metal cations (as described in U.S. Patent 5,011,912), and has been deposited with the American Type Culture Collection under accession no HB 9259. Expression systems useful for producing recombinant proteins fused to the FLAG® peptide, as well as monoclonal antibodies that bind the octapeptide and are useful in
20 purifying the recombinant proteins, are available from Eastman Kodak Company, Scientific Imaging Systems, New Haven, Connecticut. In one embodiment, the FLAG® peptide is fused to the C-terminal end of a flk-1bp polypeptide.

The present invention further includes flk-1bp polypeptides with or without associated native-pattern glycosylation. flk-1bp expressed in yeast or mammalian
25 expression systems may be similar to or significantly different from a native flk-1bp polypeptide in molecular weight and glycosylation pattern, depending upon the choice of expression system. Expression of flk-1bp polypeptides in bacterial expression systems, such as *E. coli*, provides non-glycosylated molecules.

Glycosylation sites in the flk-1bp extracellular domain can be modified to preclude
30 glycosylation while allowing expression of a homogeneous, reduced carbohydrate analog using yeast or mammalian expression systems. N-glycosylation sites in eukaryotic polypeptides are characterized by an amino acid triplet Asn-X-Y, wherein X is any amino acid except Pro and Y is Ser or Thr. Appropriate modifications to the nucleotide sequence encoding this triplet will result in substitutions, additions or deletions that prevent
35 attachment of carbohydrate residues at the Asn side chain. Known procedures for inactivating N-glycosylation sites in proteins include those described in U.S. Patent

5,071,972 and EP 276,846. Three potential N-glycosylation sites are found in SEQ ID NO:2, at positions 155-157, 185-187, and 220-222.

In another example, sequences encoding Cys residues that are not essential for biological activity can be altered to cause the Cys residues to be deleted or replaced with other amino acids, preventing formation of incorrect intramolecular disulfide bridges upon renaturation. Other variants are prepared by modification of adjacent dibasic amino acid residues to enhance expression in yeast systems in which KEX2 protease activity is present. EP 212,914 discloses the use of site-specific mutagenesis to inactivate KEX2 protease processing sites in a protein. KEX2 protease processing sites are inactivated by deleting, adding or substituting residues to alter Arg-Arg, Arg-Lys, and Lys-Arg pairs to eliminate the occurrence of these adjacent basic residues. Lys-Lys pairings are considerably less susceptible to KEX2 cleavage, and conversion of Arg-Lys or Lys-Arg to Lys-Lys represents a conservative and preferred approach to inactivating KEX2 sites. Potential KEX2 protease processing sites are found at positions 66-67, 107-108, 206-207, 324-325, 365-366, and 395-395 in SEQ ID NO:2.

flk-1bp polypeptides of the present invention, which include fragments, variants, and derivatives of native flk-1bp proteins, may be tested for the ability to bind a particular receptor using conventional assay techniques. Procedures for conducting binding assays, including competition binding assays, are well known.

A flk-1bp may be labeled with a detectable reagent (e.g., radioiodinated). Cells expressing flk-1 are contacted with the labeled flk-1bp, then washed to remove unbound reagents. Cell-associated radioactivity indicates binding of flk-1bp to the cells.

As illustrated in example 2, a flk-1bp may be tested for the ability to compete with VEGF for binding to flk-1 (i.e., a competitive binding assay). Likewise, biological activity of a flk-1bp variant, derivative, or fragment may be assessed by assaying for the variant's ability to compete with VEGF, or with a native flk-1bp, for binding to flk-1.

Reagents that may be employed in competitive binding assays include radiolabeled flk-1bp and intact flk-1-expressing cells. For example, radiolabeled native flk-1bp can be used to compete with a flk-1bp variant for binding to cell surface flk-1. Instead of intact cells, one could substitute a soluble flk-1/Fc fusion protein bound to a solid phase through the interaction of Protein A or Protein G with the Fc moiety. Chromatography columns that contain Protein A and Protein G include those available from Pharmacia Biotech, Inc., Piscataway, NJ. Another type of competitive binding assay utilizes radiolabeled soluble flk-1 and immobilized flk-1bp (e.g., an immobilized flk-1bp/Fc fusion protein).

Qualitative results can be obtained by competitive autoradiographic plate binding assays, or Scatchard plots may be utilized to generate quantitative results. Affinity calculations

(Scatchard, *Ann. N.Y. Acad. Sci.* 51:660, 1949) may be generated on RS/1 (BBN Software, Boston, MA) run on a Microvax computer.

As another alternative, binding may be detected by using a biosensor unit. One example of such a procedure involves immobilizing a flk-1/Fc fusion protein on the chip of a biosensor unit, as follows. Goat anti-human IgG directed against the Fc region (Jackson Immunoresearch Laboratories, Inc., West Grove, Pennsylvania) was chemically bound to the chip of a BIAcore Processing Unit (Pharmacia Biosensor) by standard techniques. A flk-1/Fc protein was then bound to the immobilized goat anti-human IgG *via* interaction of the antibody with the Fc moiety of the fusion protein. Culture medium from cells expressing a flk-1bp then is allowed to flow across the chip. Binding of a protein to the immobilized flk-1/Fc is indicated by a significant resonance shift on the Biosensor. The presence of a protein that binds flk-1bp in the culture medium is thus confirmed.

Oligomers

The present invention encompasses flk-1bp polypeptides in the form of oligomers, such as dimers, trimers, or higher oligomers. Oligomers may be formed by disulfide bonds between cysteine residues on different flk-1bp polypeptides, or by non-covalent interactions between flk-1bp polypeptide chains, for example. In other embodiments, oligomers comprise from two to four flk-1bp polypeptides joined *via* covalent or non-covalent interactions between peptide moieties fused to the flk-1bp polypeptides. Such peptides may be peptide linkers (spacers), or peptides that have the property of promoting oligomerization. Leucine zippers and certain polypeptides derived from antibodies are among the peptides that can promote oligomerization of flk-1bp polypeptides attached thereto, as described in more detail below.

Preparation of fusion proteins comprising heterologous polypeptides fused to various portions of antibody-derived polypeptides (including the Fc domain) has been described, e.g., by Ashkenazi et al. (*PNAS USA* 88:10535, 1991); Byrn et al. (*Nature* 344:667, 1990); and Hollenbaugh and Aruffo ("Construction of Immunoglobulin Fusion Proteins", in *Current Protocols in Immunology, Supplement 4*, pages 10.19.1 - 10.19.11, 1992), hereby incorporated by reference. In one embodiment of the invention, a flk-1bp dimer is created by fusing flk-1bp to an Fc region polypeptide derived from an antibody. The term "Fc polypeptide" includes native and mutein forms, as well as truncated Fc polypeptides containing the hinge region that promotes dimerization.

A gene fusion encoding the flk-1bp/Fc fusion protein is inserted into an appropriate expression vector. The flk-1bp/Fc fusion proteins are allowed to assemble much like antibody molecules, whereupon interchain disulfide bonds form between the Fc polypeptides, yielding divalent flk-1bp. In other embodiments, flk-1bp may be substituted

for the variable portion of an antibody heavy or light chain. If fusion proteins are made with both heavy and light chains of an antibody, it is possible to form an flk-1bp oligomer with as many as four flk-1bp polypeptides.

One suitable Fc polypeptide is the native Fc region polypeptide derived from a human IgG1, which is described in PCT application WO 93/10151, hereby incorporated by reference. Another useful Fc polypeptide is the Fc mutein described in U.S. Patent 5,457,035 and in Baum et al., *EMBO J.* 13:3992, 1994. The amino acid sequence of the mutein is identical to that of the native Fc sequence presented in WO 93/10151, except that amino acid 19 has been changed from Leu to Ala, amino acid 20 has been changed from Leu to Glu, and amino acid 22 has been changed from Gly to Ala. This mutein Fc exhibits reduced affinity for immunoglobulin receptors.

Alternatively, oligomeric flk-1bp may comprise two or more flk-1bp polypeptides joined through peptide linkers. Examples include those peptide linkers described in United States Patent 5,073,627 (hereby incorporated by reference). Fusion proteins comprising multiple flk-1bp polypeptides separated by peptide linkers may be produced using conventional recombinant DNA technology.

Another method for preparing oligomeric flk-1bp polypeptides involves use of a leucine zipper. Leucine zipper domains are peptides that promote oligomerization of the proteins in which they are found. Leucine zippers were originally identified in several DNA-binding proteins (Landschulz et al., *Science* 240:1759, 1988), and have since been found in a variety of different proteins. Among the known leucine zippers are naturally occurring peptides and derivatives thereof that dimerize or trimerize. Examples of leucine zipper domains suitable for producing oligomeric flk-1bp proteins are those described in PCT application WO 94/10308, hereby incorporated by reference. Recombinant fusion proteins comprising a flk-1bp polypeptide fused to a peptide that dimerizes or trimerizes in solution are expressed in suitable host cells, and the resulting oligomeric flk-1bp is recovered from the culture supernatant.

In one embodiment, a leucine zipper moiety that preferentially forms trimers is employed. One example is a leucine zipper derived from lung surfactant protein D (SPD), as described in Hoppe et al. (*FEBS Letters* 344:191, 1994). Other peptides derived from naturally occurring trimeric proteins may be employed in preparing trimeric flk-1bp.

Expression Systems

The present invention provides recombinant expression vectors for expression of flk-1bp, and host cells transformed with the expression vectors. Any suitable expression system may be employed. The vectors include a DNA encoding a flk-1bp polypeptide, operably linked to suitable transcriptional or translational regulatory nucleotide sequences,

such as those derived from a mammalian, microbial, viral, or insect gene. Examples of regulatory sequences include transcriptional promoters, operators, or enhancers, an mRNA ribosomal binding site, and appropriate sequences which control transcription and translation initiation and termination. Nucleotide sequences are operably linked when the regulatory sequence functionally relates to the flk-1bp DNA sequence. Thus, a promoter nucleotide sequence is operably linked to an flk-1bp DNA sequence if the promoter nucleotide sequence controls the transcription of the flk-1bp DNA sequence. An origin of replication that confers the ability to replicate in the desired host cells, and a selection gene by which transformants are identified, are generally incorporated into the expression vector.

Advantageously, a sequence encoding an appropriate signal peptide, either native or heterologous, is incorporated into the expression vectors. A DNA sequence for a signal peptide (secretory leader) may be fused in frame to the flk-1bp sequence so that the flk-1bp is initially translated as a fusion protein comprising the signal peptide. A signal peptide that is functional in the intended host cells promotes extracellular secretion of the flk-1bp polypeptide. The signal peptide is cleaved from the flk-1bp polypeptide upon secretion of flk-1bp from the cell. Heterologous signal or leader peptides may be employed when such peptides promote secretion of higher levels of flk-1bp from the particular host cells employed.

Suitable host cells for expression of flk-1bp polypeptides include prokaryotes, yeast or higher eukaryotic cells. Appropriate cloning and expression vectors for use with bacterial, fungal, yeast, and mammalian cellular hosts are described, for example, in Pouwels et al. *Cloning Vectors: A Laboratory Manual*, Elsevier, New York, (1985). Cell-free translation systems could also be employed to produce flk-1bp polypeptides using RNAs derived from DNA constructs disclosed herein.

Prokaryotes include gram negative or gram positive organisms, for example, *E. coli* or *Bacilli*. Suitable prokaryotic host cells for transformation include, for example, *E. coli*, *Bacillus subtilis*, *Salmonella typhimurium*, and various other species within the genera *Pseudomonas*, *Streptomyces*, and *Staphylococcus*. In a prokaryotic host cell, such as *E. coli*, a flk-1bp polypeptide may include an N-terminal methionine residue to facilitate expression of the recombinant polypeptide in the prokaryotic host cell. The N-terminal Met may be cleaved from the expressed recombinant flk-1bp polypeptide.

Expression vectors for use in prokaryotic host cells generally comprise one or more phenotypic selectable marker genes. A phenotypic selectable marker gene is, for example, a gene encoding a protein that confers antibiotic resistance or that supplies an autotrophic requirement. Examples of useful expression vectors for prokaryotic host cells include those derived from commercially available plasmids such as the cloning vector pBR322

(ATCC 37017). pBR322 contains genes for ampicillin and tetracycline resistance and thus provides simple means for identifying transformed cells. An appropriate promoter and a flk-1bp DNA sequence are inserted into the pBR322 vector. Other commercially available vectors include, for example, pKK223-3 (Pharmacia Fine Chemicals, Uppsala, Sweden) and pGEM1 (Promega Biotec, Madison, WI, USA).

Promoter sequences commonly used for recombinant prokaryotic host cell expression vectors include β -lactamase (penicillinase), lactose promoter system (Chang et al., *Nature* 275:615, 1978; and Goeddel et al., *Nature* 281:544, 1979), tryptophan (trp) promoter system (Goeddel et al., *Nucl. Acids Res.* 8:4057, 1980; and EP-A-36776) and tac promoter (Maniatis, *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratory, p. 412, 1982). A particularly useful prokaryotic host cell expression system employs a phage λ P_L promoter and a cI857ts thermolabile repressor sequence. Plasmid vectors available from the American Type Culture Collection which incorporate derivatives of the λ P_L promoter include plasmid pHUB2 (resident in *E. coli* strain JMB9, ATCC 37092) and pPLc28 (resident in *E. coli* RR1, ATCC 53082).

flk-1bp alternatively may be expressed in yeast host cells, preferably from the *Saccharomyces* genus (e.g., *S. cerevisiae*). Other genera of yeast, such as *Pichia* or *Kluyveromyces*, may also be employed. Yeast vectors will often contain an origin of replication sequence from a 2 μ yeast plasmid, an autonomously replicating sequence (ARS), a promoter region, sequences for polyadenylation, sequences for transcription termination, and a selectable marker gene.

Suitable promoter sequences for yeast vectors include, among others, promoters for metallothionein, 3-phosphoglycerate kinase (Hitzeman et al., *J. Biol. Chem.* 255:2073, 1980) or other glycolytic enzymes (Hess et al., *J. Adv. Enzyme Reg.* 7:149, 1968; and Holland et al., *Biochem.* 17:4900, 1978), such as enolase, glyceraldehyde-3-phosphate dehydrogenase, hexokinase, pyruvate decarboxylase, phosphofructokinase, glucose-6-phosphate isomerase, 3-phosphoglycerate mutase, pyruvate kinase, triosephosphate isomerase, phospho-glucose isomerase, and glucokinase. Other suitable vectors and promoters for use in yeast expression are further described in Hitzeman, EPA-73,657. Another alternative is the glucose-repressible ADH2 promoter described by Russell et al. (*J. Biol. Chem.* 258:2674, 1982) and Beier et al. (*Nature* 300:724, 1982). Shuttle vectors replicable in both yeast and *E. coli* may be constructed by inserting DNA sequences from pBR322 for selection and replication in *E. coli* (Amp^r gene and origin of replication) into the above-described yeast vectors.

The yeast α -factor leader sequence may be employed to direct secretion of the flk-1bp polypeptide. The α -factor leader sequence is often inserted between the promoter sequence and the structural gene sequence. See, e.g., Kurjan et al., *Cell* 30:933, 1982 and

Bitter et al., *Proc. Natl. Acad. Sci. USA* 81:5330, 1984. Other leader sequences suitable for facilitating secretion of recombinant polypeptides from yeast hosts are known to those of skill in the art. A leader sequence may be modified near its 3' end to contain one or more restriction sites. This will facilitate fusion of the leader sequence to the structural gene.

Yeast transformation protocols are known to those of skill in the art. One such protocol is described by Hinnen et al., *Proc. Natl. Acad. Sci. USA* 75:1929, 1978. The Hinnen et al. protocol selects for Trp⁺ transformants in a selective medium, wherein the selective medium consists of 0.67% yeast nitrogen base, 0.5% casamino acids, 2% glucose, 10 µg/ml adenine and 20 µg/ml uracil.

Yeast host cells transformed by vectors containing an ADH2 promoter sequence may be grown for inducing expression in a "rich" medium. An example of a rich medium is one consisting of 1% yeast extract, 2% peptone, and 1% glucose supplemented with 80 µg/ml adenine and 80 µg/ml uracil. Derepression of the ADH2 promoter occurs when glucose is exhausted from the medium.

Mammalian or insect host cell culture systems could also be employed to express recombinant flk-1bp polypeptides. Baculovirus systems for production of heterologous proteins in insect cells are reviewed by Luckow and Summers, *BioTechnology* 6:47 (1988). Established cell lines of mammalian origin also may be employed. Examples of suitable mammalian host cell lines include the COS-7 line of monkey kidney cells (ATCC CRL 1651) (Gluzman et al., *Cell* 23:175, 1981), L cells, C127 cells, 3T3 cells (ATCC CCL 163), Chinese hamster ovary (CHO) cells, HeLa cells, and BHK (ATCC CRL 10) cell lines, and the CVI/EBNA cell line derived from the African green monkey kidney cell line CVI (ATCC CCL 70) as described by McMahan et al. (*EMBO J.* 10: 2821, 1991).

Transcriptional and translational control sequences for mammalian host cell expression vectors may be excised from viral genomes. Commonly used promoter sequences and enhancer sequences are derived from Polyoma virus, Adenovirus 2, Simian Virus 40 (SV40), and human cytomegalovirus. DNA sequences derived from the SV40 viral genome, for example, SV40 origin, early and late promoter, enhancer, splice, and polyadenylation sites may be used to provide other genetic elements for expression of a structural gene sequence in a mammalian host cell. Viral early and late promoters are particularly useful because both are easily obtained from a viral genome as a fragment which may also contain a viral origin of replication (Fiers et al., *Nature* 273:113, 1978). Smaller or larger SV40 fragments may also be used, provided the approximately 250 bp sequence extending from the *Hind* III site toward the *Bgl* I site located in the SV40 viral origin of replication site is included.

Expression vectors for use in mammalian host cells can be constructed as disclosed by Okayama and Berg (*Mol. Cell. Biol.* 3:280, 1983), for example. A useful system for stable high level expression of mammalian cDNAs in C127 murine mammary epithelial cells can be constructed substantially as described by Cosman et al. (*Mol. Immunol.* 23:935, 1986). A high expression vector, PMLSV N1/N4, described by Cosman et al., *Nature* 312:768, 1984 has been deposited as ATCC 39890. Additional examples of mammalian expression vectors are described in EP-A-0367566, and in WO 91/18982. The vectors may be derived from retroviruses.

If desired, a heterologous signal or leader peptide may be substituted for the native flk-1bp signal peptide. Signal or leader peptides are chosen according to the particular host cell that is to be employed.

Examples of alternative signal peptides that are functional in mammalian host cells include the signal sequence for interleukin-7 (IL-7) described in United States Patent 4,965,195; the signal sequence for interleukin-2 receptor described in Cosman et al., *Nature* 312:768 (1984); the interleukin-4 receptor signal peptide described in EP 367,566; the type I interleukin-1 receptor signal peptide described in U.S. Patent 4,968,607; and the type II interleukin-1 receptor signal peptide described in EP 460,846. Synthetic signal peptides, which are not naturally occurring but can be generated through recombinant DNA technology, are yet another alternative.

Purified flk-1bp Protein

The present invention provides purified flk-1bp proteins, which may be produced by recombinant expression systems as described above or purified from naturally occurring cells. The desired degree of purity may depend on the intended use of the protein. A relatively high degree of purity is desired when the protein is to be administered *in vivo*, for example. Advantageously, flk-1bp polypeptides are purified such that no protein bands corresponding to other proteins are detectable by SDS-polyacrylamide gel electrophoresis (SDS-PAGE). It will be recognized by one skilled in the pertinent field that multiple bands corresponding to flk-1bp protein may be detected by SDS-PAGE, due to differential glycosylation, variations in post-translational processing, and the like, as discussed above. A preparation of flk-1bp protein is considered to be purified as long as no bands corresponding to different (non-flk-1bp) proteins are visualized. flk-1bp most preferably is purified to substantial homogeneity, as indicated by a single protein band upon analysis by SDS-PAGE. The protein band may be visualized by silver staining, Coomassie blue staining, or (if the protein is radiolabeled) by autoradiography.

One process for producing the flk-1bp protein comprises culturing a host cell transformed with an expression vector comprising a DNA sequence that encodes flk-1bp

under conditions such that flk-1bp is expressed. The flk-1bp protein is then recovered from the culture medium. As the skilled artisan will recognize, procedures for purifying the recombinant flk-1bp will vary according to such factors as the type of host cells employed.

5 For example, the culture medium first may be concentrated using a commercially available protein concentration filter, for example, an Amicon or Millipore Pellicon ultrafiltration unit. Following the concentration step, the concentrate can be applied to a purification matrix such as a gel filtration medium. Alternatively, an anion exchange resin can be employed, for example, a matrix or substrate having pendant diethylaminoethyl
10 (DEAE) groups. The matrices can be acrylamide, agarose, dextran, cellulose or other types commonly employed in protein purification. Alternatively, a cation exchange step can be employed. Suitable cation exchangers include various insoluble matrices comprising sulfopropyl or carboxymethyl groups. Sulfopropyl groups are preferred. Finally, one or more reversed-phase high performance liquid chromatography (RP-HPLC) steps
15 employing hydrophobic RP-HPLC media, (e.g., silica gel having pendant methyl or other aliphatic groups) can be employed to further purify flk-1bp. Some or all of the foregoing purification steps, in various combinations, can be employed to provide a purified flk-1bp protein.

 Recombinant protein produced in bacterial culture may be isolated by initial
20 disruption of the host cells, centrifugation, extraction from cell pellets if an insoluble polypeptide, or from the supernatant fluid if a soluble polypeptide, followed by one or more concentration, salting-out, ion exchange, affinity purification or size exclusion chromatography steps. Finally, RP-HPLC can be employed for final purification steps. Microbial cells can be disrupted by any convenient method, including freeze-thaw cycling,
25 sonication, mechanical disruption, or use of cell lysing agents.

 Transformed yeast host cells may be employed to express flk-1bp, preferably as a secreted polypeptide to simplify purification. Secreted recombinant polypeptide from a yeast host cell fermentation can be purified by methods analogous to those disclosed by Urdal et al. (*J. Chromatog.* 296:171, 1984). Urdal et al. describe two sequential,
30 reversed-phase HPLC steps for purification of recombinant human IL-2 on a preparative HPLC column.

 Alternatively, flk-1bp polypeptides can be purified by immunoaffinity chromatography. An affinity column containing an immobilized antibody that binds flk-1bp may be prepared by conventional procedures and employed in purifying flk-1bp.
35 Example 3 describes a procedure for generating monoclonal antibodies directed against flk-1bp.

Properties and Uses of flk-1bp

The flk-1bp of the present invention binds to the receptor known as human flk-1 or KDR (Terman et al., *Oncogene* 6:1677-83, 1991; Terman et al., *Biochem. Biophys. Res. Comm.* 187:1579-86, 1992). As demonstrated in example 2, human flk-1bp competes
5 with VEGF for binding to human flk-1. A number of uses flow from this property of flk-1bp.

flk-1 is expressed on vascular endothelial cells. flk-1bp polypeptides may be used to purify vascular endothelial cells, or any cell type to which flk-1bp binds. In one embodiment, the cells are purified by affinity chromatography, using a chromatography
10 matrix having flk-1bp bound thereto. The flk-1bp attached to the chromatography matrix may be a full length protein, a flk-1bp fragment, a flk-1bp-containing fusion protein, or other suitable flk-1bp polypeptide described herein. In one embodiment, a flk-1bp/Fc fusion protein is bound to a Protein A or Protein G column through interaction of the Fc moiety with the Protein A or Protein G. Alternatively, flk-1bp may be used in isolating
15 vascular endothelial cells by flow cytometry.

The flk-1bp also finds use as a protein purification reagent. Human flk-1 may be purified by affinity chromatography, for example, using a chromatography matrix having flk-1bp bound thereto.

Preparations of vascular endothelial cells and flk-1 are useful to those studying the
20 roles of such cells and receptor tyrosine kinases in vasculogenesis and angiogenesis (formation and sprouting of blood vessels, respectively) (Millauer et al., *supra*, and Mustonen et al., *supra*). The disclosure herein of the flk-1bp of the present invention allows investigation of the role that this novel protein may play in vasculogenesis or angiogenesis.

Angiogenesis is essential for a number of normal processes (such as embryonic development, somatic growth, cyclical growth of endometrium, and wound healing), yet has been implicated in certain pathological conditions as well (e.g., rheumatoid arthritis and the growth and metastasis of solid tumors). See Ferrara et al. (*J. Cell. Biochem.* 47:211, 1991; and Folkman, J. (*Nature Med.*, 1:27-31, 1995), hereby incorporated by reference.

The flk-1bp of the present invention may be used in developing treatments for any disorder mediated (directly or indirectly) by defective or insufficient amounts of flk-1bp. A therapeutically effective amount of purified flk-1bp protein is administered to a patient afflicted with such a disorder. Alternatively, flk-1bp DNA sequences may be employed in developing a gene therapy approach to treating such disorders. Disclosure herein of native
35 flk-1bp nucleotide sequences permits the detection of defective flk-1bp genes, and the replacement thereof with normal flk-1bp-encoding genes. Defective genes may be detected in *in vitro* diagnostic assays, and by comparison of the native flk-1bp nucleotide sequence

disclosed herein with that of an flk-1bp gene derived from a person suspected of harboring a defect in this gene.

Polypeptides of the present invention also may be employed as carriers, for delivering diagnostic or therapeutic agents to any cells to which flk-1bp binds. As discussed above, such cells include but are not limited to cells bearing flk-1. Flk-1bp can be used to deliver diagnostic or therapeutic agents to these cells in *in vitro* or *in vivo* procedures. Such cells are contacted with a conjugate comprising a diagnostic or therapeutic agent attached to a flk-1bp polypeptide. The flk-1bp binds to the target cells, thus allowing detection thereof (in the case of diagnostic agents) or treatment thereof (with therapeutic agents).

Flk-1bp/diagnostic agent conjugates may be employed to detect the presence of flk-1⁺ cells *in vitro* or *in vivo*. Conjugates containing a cytotoxic agent attached to flk-1bp may be used to kill flk-1⁺ cells. In one embodiment, such conjugates containing cytotoxic agents are employed to kill flk-1⁺ cells involved in angiogenesis.

Diagnostic and therapeutic agents that may be attached to a flk-1bp polypeptide include, but are not limited to, drugs, toxins, radionuclides, chromophores, enzymes that catalyze a colorimetric or fluorometric reaction, and the like, with the particular agent being chosen according to the intended application. Among the toxins are ricin, abrin, saporin toxin, diphtheria toxin, *Pseudomonas aeruginosa* exotoxin A, ribosomal inactivating proteins, mycotoxins such as trichothecenes, and derivatives and fragments (e.g., single chains) thereof. Radionuclides suitable for diagnostic use include, but are not limited to, ¹²³I, ¹³¹I, ^{99m}Tc, ¹¹¹In, and ⁷⁶Br. Radionuclides suitable for therapeutic use include, but are not limited to, ¹³¹I, ²¹¹At, ⁷⁷Br, ¹⁸⁶Re, ¹⁸⁸Re, ²¹²Pb, ²¹²Bi, ¹⁰⁹Pd, ⁶⁴Cu, and ⁶⁷Cu.

Such agents may be attached to the flk-1bp by any suitable conventional procedure. Being a protein, flk-1bp comprises functional groups on amino acid side chains that can be reacted with functional groups on a desired agent to form covalent bonds, for example. The agent may be covalently linked to flk-1bp via an amide bond, hindered disulfide bond, acid-cleavable linkage, and the like, which are among the conventional linkages chosen according to such factors as the structure of the desired agent. Alternatively, flk-1bp or the agent to be linked thereto may be derivatized to generate or attach a desired reactive functional group. The derivatization may involve attachment of one of the bifunctional coupling reagents available for linking various molecules to proteins (Pierce Chemical Company, Rockford, Illinois). A number of techniques for radiolabeling proteins are known. One such method involves use of commercially available reagents (Pierce Chemical Company) to radioiodinate a flk-1bp polypeptide. Radionuclide metals may be attached to flk-1bp by using a suitable bifunctional chelating agent, examples of which are described in U.S. patents 4,897,255 and 4,965,392.

Conjugates comprising flk-1bp and a suitable diagnostic or therapeutic agent (preferably covalently linked) are thus prepared. The conjugates are administered or otherwise employed in an amount appropriate for the particular indication.

5 The present invention provides pharmaceutical compositions comprising purified flk-1bp and a physiologically acceptable carrier, diluent, or excipient. Such compositions may comprise flk-1bp in any form described herein, e.g., full length native flk-1bp, or fragments, variants, oligomers, derivatives, or conjugates thereof. Components that are commonly employed in pharmaceutical formulations include those described in *Remington's Pharmaceutical Sciences and U.S. Pharmacopoeia: National Formulary*,
10 Mack Publishing Company, Easton, Pennsylvania, 1984. Suitable carriers, diluents, and excipients are nontoxic to recipients at the dosages and concentrations employed. Such compositions may comprise buffers, antioxidants such as ascorbic acid, low molecular weight (less than about 10 residues) polypeptides, proteins, amino acids, carbohydrates including glucose, sucrose or dextrans, chelating agents such as EDTA, glutathione and
15 other stabilizers and excipients commonly employed in pharmaceutical compositions. Neutral buffered saline or saline mixed with conspecific serum albumin are among the appropriate diluents. The composition may be formulated as a lyophilizate using appropriate excipient solutions (e.g. sucrose) as diluents.

For therapeutic use, purified proteins of the present invention are administered to a
20 patient, preferably a human, for treatment in a manner appropriate to the indication. Thus, for example, the pharmaceutical compositions can be administered locally, by intravenous injection, continuous infusion, sustained release from implants, or other suitable technique. Appropriate dosages and the frequency of administration will depend, of course, on such factors as the nature and severity of the indication being treated, the desired response, the
25 condition of the patient and so forth.

The flk-1bp protein employed in the pharmaceutical compositions preferably is purified such that the flk-1bp protein is substantially free of other proteins of natural or endogenous origin, desirably containing less than about 1% by mass of protein contaminants residual of production processes. Such compositions, however, can contain
30 other proteins added as stabilizers, carriers, excipients or co-therapeutics.

The flk-1bp of the present invention also finds use in various *in vitro* assays. In one such assay, a molecule is tested for the ability to compete with flk-1bp for binding to a given receptor. Other assays identify molecules that modulate angiogenesis. Such molecules may be proteins or small peptides, or may be non-proteinaceous in nature, and
35 include naturally occurring or synthetic molecules.

Flk-1bp may be employed in screening assays to identify antagonists, i.e., molecules capable of inhibiting a biological activity of flk-1bp. In one embodiment, the assay identifies molecules that inhibit flk-1bp-mediated angiogenesis.

One method for identifying antagonists of flk-1bp comprises contacting cells
5 expressing a receptor that binds flk-1bp with flk-1bp in the presence of a candidate antagonist, and analyzing whether the candidate antagonist inhibits a biological effect induced by the binding of flk-1bp to the cells. One of the above-described diagnostic (detectable) agents may be attached to the flk-1bp. In one embodiment, the receptor is flk-1. Suitable cells include, but are not limited to, flk-1⁺ endothelial cells, e.g., flk-1⁺ vascular
10 endothelial cells. In particular embodiments, the biological effect is proliferation of the cells or phosphorylation of the receptor.

Inhibitors (antagonists) of flk-1bp that are identified in such screening assays are provided, along with compositions containing the inhibitors. The thus-identified antagonists may be employed to inhibit a biological activity of flk-1bp, e.g., may be
15 administered *in vivo* to treat conditions mediated (directly or indirectly) by flk-1bp. The antagonists find use in inhibiting angiogenesis, including the angiogenesis associated with growth of solid tumors.

The flk-1bp-encoding DNAs disclosed herein find use in the production of flk-1bp polypeptides, as discussed above. Fragments of the flk-1bp nucleotide sequences are also
20 useful. In one embodiment, such fragments comprise at least about 17 consecutive nucleotides, more preferably at least 30 consecutive nucleotides, of the flk-1bp DNA disclosed herein. DNA and RNA complements of said fragments are provided herein, along with both single-stranded and double-stranded forms of the flk-1bp DNA of SEQ ID NO:1.

Among the uses of such flk-1bp nucleic acid fragments are use as a probe or as
25 primers in a polymerase chain reaction (PCR). As one example, such a probe finds use in detecting the presence of flk-1bp nucleic acids in *in vitro* assays and in such procedures as Northern and Southern blots. Cell types expressing flk-1bp can be identified as well. Such procedures are well known, and the skilled artisan can choose a probe of suitable
30 length, depending on the particular intended application. For PCR, 5' and 3' primers corresponding to the termini of a desired flk-1bp DNA are employed in isolating and amplifying the DNA, using conventional techniques.

Other useful fragments of flk-1bp nucleic acids are antisense or sense
oligonucleotides comprising a single-stranded nucleic acid sequence (either RNA or DNA)
35 capable of binding to target flk-1bp mRNA (sense) or flk-1bp DNA (antisense) sequences. Such a fragment generally comprises at least about 14 nucleotides, preferably from about 14 to about 30 nucleotides. The ability to create an antisense or a sense oligonucleotide,

based upon a cDNA sequence for a given protein is described in, for example, Stein and Cohen, *Cancer Res.* 48:2659, 1988 and van der Krol et al., *BioTechniques* 6:958, 1988.

Binding of antisense or sense oligonucleotides to target nucleic acid sequences results in the formation of duplexes that block translation (RNA) or transcription (DNA) by one of several means, including enhanced degradation of the duplexes, premature termination of transcription or translation, or by other means. The antisense oligonucleotides thus may be used to block expression of flk-1bp proteins.

Antisense or sense oligonucleotides further comprise oligonucleotides having modified sugar-phosphodiester backbones (or other sugar linkages, such as those described in WO 91/06629) and wherein such sugar linkages are resistant to endogenous nucleases. Such oligonucleotides with resistant sugar linkages are stable *in vivo* (i.e., capable of resisting enzymatic degradation) but retain sequence specificity to be able to bind to target nucleotide sequences. Other examples of sense or antisense oligonucleotides include those oligonucleotides which are covalently linked to organic moieties, such as those described in WO 90/10448, and other moieties that increases affinity of the oligonucleotide for a target nucleic acid sequence, such as poly-(L-lysine). Further still, intercalating agents, such as ellipticine, and alkylating agents or metal complexes may be attached to sense or antisense oligonucleotides to modify binding specificities of the antisense or sense oligonucleotide for the target nucleotide sequence.

Antisense or sense oligonucleotides may be introduced into a cell containing the target nucleic acid sequence by any gene transfer method, including, for example, CaPO₄-mediated DNA transfection, electroporation, or other gene transfer vectors such as Epstein-Barr virus. Antisense or sense oligonucleotides are preferably introduced into a cell containing the target nucleic acid sequence by insertion of the antisense or sense oligonucleotide into a suitable retroviral vector, then contacting the cell with the retrovirus vector containing the inserted sequence, either *in vivo* or *ex vivo*. Suitable retroviral vectors include, but are not limited to, the murine retrovirus M-MuLV, N2 (a retrovirus derived from M-MuLV), or the double copy vectors designated DCT5A, DCT5B and DCT5C (see PCT Application US 90/02656). Alternatively, other promotor sequences may be used to express the oligonucleotide.

Sense or antisense oligonucleotides may also be introduced into a cell containing the target nucleotide sequence by formation of a conjugate with a ligand binding molecule, as described in WO 91/04753. Suitable ligand binding molecules include, but are not limited to, cell surface receptors, growth factors, other cytokines, or other ligands that bind to cell surface receptors. Preferably, conjugation of the ligand binding molecule does not substantially interfere with the ability of the ligand binding molecule to bind to its

corresponding molecule or receptor, or block entry of the sense or antisense oligonucleotide or its conjugated version into the cell.

- Alternatively, a sense or an antisense oligonucleotide may be introduced into a cell containing the target nucleic acid sequence by formation of an oligonucleotide-lipid complex, as described in WO 90/10448. The sense or antisense oligonucleotide-lipid complex is preferably dissociated within the cell by an endogenous lipase.

Antibodies Immunoreactive with flk-1bp

- The flk-1bp protein of the present invention, or immunogenic fragments thereof, may be employed in generating antibodies. The present invention thus provides antibodies that specifically bind flk-1bp, i.e., the antibodies bind to flk-1bp *via* the antigen-binding sites of the antibody (as opposed to non-specific binding).

- Polyclonal and monoclonal antibodies directed against flk-1bp may be prepared by conventional techniques. See, for example, *Monoclonal Antibodies, Hybridomas: A New Dimension in Biological Analyses*, Kennet et al. (eds.), Plenum Press, New York (1980); and *Antibodies: A Laboratory Manual*, Harlow and Land (eds.), Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY, (1988). Production of monoclonal antibodies that are immunoreactive with flk-1bp is further illustrated in example 3 below.

- Antigen-binding fragments of such antibodies, which may be produced by conventional techniques, are also encompassed by the present invention. Examples of such fragments include, but are not limited to, Fab, F(ab'), and F(ab')₂ fragments. Antibody fragments and derivatives produced by genetic engineering techniques are also provided.

- The monoclonal antibodies of the present invention include chimeric antibodies, e.g., humanized versions of murine monoclonal antibodies. Such humanized antibodies may be prepared by known techniques, and offer the advantage of reduced immunogenicity when the antibodies are administered to humans. In one embodiment, a humanized monoclonal antibody comprises the variable region of a murine antibody (or just the antigen binding site thereof) and a constant region derived from a human antibody. Alternatively, a humanized antibody fragment may comprise the antigen binding site of a murine monoclonal antibody and a variable region fragment (lacking the antigen-binding site) derived from a human antibody. Procedures for the production of chimeric and further engineered monoclonal antibodies include those described in Riechmann et al. (*Nature* 332:323, 1988), Liu et al. (*PNAS* 84:3439, 1987), Larrick et al. (*Bio/Technology* 7:934, 1989), and Winter and Harris (*TIPS* 14:139, May, 1993).

- Among the uses of the antibodies is use in assays to detect the presence of flk-1bp polypeptides, either *in vitro* or *in vivo*. The antibodies find further use in purifying flk-1bp by affinity chromatography.

Certain of the antibodies may additionally be able to block binding of flk-1bp to target cells and to inhibit a biological activity of flk-1bp *in vivo*. A therapeutic method involves *in vivo* administration of such an antibody in an amount effective in inhibiting an flk-1bp-mediated biological activity. Disorders mediated or exacerbated by flk-1bp, directly or indirectly, are thus treated. In one embodiment, such an antibody is administered to a mammal to inhibit angiogenesis.

Pharmaceutical compositions comprising an antibody that specifically binds flk-1bp, and a suitable, diluent, excipient, or carrier, are provided herein. Suitable components of such compositions are as described above for the compositions containing flk-1bp proteins.

The following examples are provided to illustrate particular embodiments of the present invention, and are not to be construed as limiting the scope of the invention.

EXAMPLE 1: Isolation of a Human flk-1bp DNA

DNA encoding a human flk-1bp protein was isolated by the following procedure. A search of a sequence databank was performed, using a vascular endothelial cell growth factor (VEGF) amino acid sequence as the query sequence. An expressed sequence tag (EST) file, GenBank accession number Z44272, was identified by the search.

Two oligonucleotides based upon sequences at the 3' and 5' ends of this 299-nucleotide EST file were synthesized for use as polymerase chain reaction (PCR) primers. PCR was conducted by conventional procedures, using DNA from a WI-26VA4 (human fibroblast cell line; see below) cDNA library in a λ gt10 vector as the template. A 299-bp DNA fragment that was isolated and amplified by the PCR corresponded to nucleotides 660 to 958 of SEQ ID NO:1.

A cDNA library derived from a human fibroblast cell line designated WI-26VA4 was prepared by isolating polyA⁺ RNA from WI-26VA4 cells cultured in the presence of pokeweed mitogen. cDNA was synthesized on the mRNA template by conventional techniques. Double stranded cDNA at least about 500bp in length was inserted into an expression vector designated pDC302. The mammalian expression vector pDC302, which also replicates in *E. coli*, is described in Mosley et al. (*Cell* 59:335-348, 1989).

The 299-bp DNA fragment isolated above was labeled with ³²P and used as a probe to screen the human fibroblast cDNA library. The nucleotide sequence of a positive clone, designated human flk-1bp clone 9C1, was determined, and is presented in SEQ ID NO:1. The amino acid sequence encoded thereby is presented in SEQ ID NO:2. This human flk-1bp is a secreted (as opposed to cell membrane bound) protein comprising an N-terminal signal peptide (amino acids -20 to -1 of SEQ ID NO:2) followed by the mature protein

(amino acids 1 to 399). The calculated molecular weight of the mature form of this protein is 44,823 daltons, and the predicted pI is 8.678.

E. coli strain DH10B cells transformed with a recombinant vector containing this flk-1bp DNA were deposited with the American Type Culture Collection on September 1, 1995, and assigned accession no. ATCC 69897. The deposit was made under the terms of the Budapest Treaty. The vector in the deposited strain is the cloning vector pBluescript®SK(-) (Stratagene Cloning Systems, LaJolla, CA). The vector was digested with EcoRI (found in the multiple cloning site of the vector), and human flk-1bp DNA that includes the entire coding region shown in SEQ ID NO:1, as well as additional 5' and 3' non-coding sequences, was ligated into the digested vector.

EXAMPLE 2: Competition Binding Assay

Human flk-1bp was expressed and tested for the ability to compete with VEGF for binding the receptor designated flk-1, as follows. A cDNA encoding amino acids -20 to 399 of SEQ ID NO:2 was inserted into the expression vector pDC302 (Mosley et al., *Cell* 59:335-348, 1989). CV-1/EBNA-1 cells were transfected with the resulting recombinant expression vector. The monkey kidney cell line CV-1/EBNA-1 (ATCC CRL 10478) was derived by transfection of the CV-1 cell line (ATCC CCL 70) with a gene encoding Epstein-Barr virus nuclear antigen-1 (EBNA-1) that constitutively expresses EBNA-1 driven from the human CMV intermediate-early enhancer/promoter, as described by McMahan et al. (*EMBO J.* 10:2821, 1991).

The transfected cells were cultured to allow secretion of the flk-1bp into the culture medium. Cells were pelleted by centrifugation, and culture supernatants were concentrated 10-fold for use in the following assay.

Human vascular endothelial cell growth factor (VEGF), which is known to bind flk-1, was radioiodinated using standard techniques. The ability of the flk-1bp-containing supernatant to compete with the radioiodinated VEGF for binding to a human flk-1/Fc fusion protein was tested. The fusion protein comprised the extracellular domain of human flk-1 fused to an Fc region polypeptide derived from an antibody. Flk-1/Fc fusion proteins may be prepared by procedures analogous to those described in Park et al. (*J. Biol. Chem.* 269:25646, 1994, at page 25647), hereby incorporated by reference.

The flk-1bp-containing supernatant inhibited about 80% of the binding of VEGF to the flk-1/Fc fusion protein. Control supernatant from CV1-EBNA cells inhibited about 20% of the VEGF binding to flk-1/Fc.

EXAMPLE 3: Antibodies that bind flk-1bp

This example illustrates the preparation of monoclonal antibodies that specifically bind flk-1bp. Suitable immunogens that may be employed in generating such antibodies include, but are not limited to, purified flk-1bp protein, an immunogenic fragment thereof, and fusion proteins containing flk-1bp polypeptides (e.g., flk-1bp/Fc fusion proteins).

Known techniques for producing monoclonal antibodies include those described in U.S. Patent 4,411,993. Briefly, mice are immunized with flk-1bp as an immunogen emulsified in complete Freund's adjuvant, and injected in amounts ranging from 10-100 µg subcutaneously or intraperitoneally. Ten to twelve days later, the immunized animals are boosted with additional flk-1bp emulsified in incomplete Freund's adjuvant. Mice are periodically boosted thereafter on a weekly to bi-weekly immunization schedule. Serum samples are periodically taken by retro-orbital bleeding or tail-tip excision for testing by dot blot assay or ELISA (Enzyme-Linked Immuno-sorbent Assay) for antibodies directed against flk-1bp.

Following detection of an appropriate antibody titer, positive animals are provided one last intravenous injection of flk-1bp in saline. Three to four days later, the animals are sacrificed, spleen cells harvested, and spleen cells are fused to a murine myeloma cell line such as NS1 or, preferably, P3x63Ag 8.653 (ATCC CRL 1580). Fusions generate hybridoma cells, which are plated in multiple microtiter plates in a HAT (hypoxanthine, aminopterin and thymidine) selective medium to inhibit proliferation of non-fused cells, myeloma hybrids, and spleen cell hybrids.

The hybridoma cells are screened by ELISA for reactivity against purified flk-1bp by adaptations of the techniques disclosed in Engvall et al. (*Immunochem.* 8:871, 1971) and in U.S. Patent 4,703,004. Positive hybridoma cells can be injected intraperitoneally into syngeneic BALB/c mice to produce ascites containing high concentrations of anti-flk-1bp monoclonal antibodies. Alternatively, hybridoma cells can be grown *in vitro* in flasks or roller bottles by various techniques. Monoclonal antibodies produced in mouse ascites can be purified by ammonium sulfate precipitation, followed by gel exclusion chromatography. Alternatively, affinity chromatography based upon binding of antibody to protein A or protein G can be used, as can affinity chromatography based upon binding to flk-1bp.

EXAMPLE 4: Northern Blot Analysis

Expression of flk-1bp in a number of different tissue types was analysed in a conventional northern blot procedure. Northern blots containing poly A⁺ RNA from a variety of adult human tissues was obtained from Clontech (Palo Alto, CA). The blots were probed with an antisense riboprobe derived from human flk-1bp DNA.

Human flk-1 bp mRNA was detected in heart, liver, skeletal muscle, pancreas, and prostate, with weaker but detectable expression in brain, placenta, lung, kidney, spleen, thymus, colon, ovaries, and small intestine. Human flk-1 bp message was not detected in peripheral blood leukocytes.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Immunex Corporation
- (ii) TITLE OF INVENTION: flk-1 Binding Protein
- (iii) NUMBER OF SEQUENCES: 5
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Immunex Corporation
 - (B) STREET: 51 University Street
 - (C) CITY: Seattle
 - (D) STATE: WA
 - (E) COUNTRY: US
 - (F) ZIP: 98101
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: Apple Power Macintosh
 - (C) OPERATING SYSTEM: Apple 7.5.2
 - (D) SOFTWARE: Microsoft Word, Version 6.0.1
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: --to be assigned--
 - (B) FILING DATE: 05-NOV-1996
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/554,374
 - (B) FILING DATE: 08-NOV-1995
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Anderson, Kathryn A.
 - (B) REGISTRATION NUMBER: 32,172
 - (C) REFERENCE/DOCKET NUMBER: 2837-WO
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (206) 587-0430
 - (B) TELEFAX: (206) 233-0644
 - (C) TELEX: 756822

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2321 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO

(vii) IMMEDIATE SOURCE:

(B) CLONE: hu flk-1 bp

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 188..1447

(ix) FEATURE:

(A) NAME/KEY: sig_peptide

(B) LOCATION: 188..247

(ix) FEATURE:

(A) NAME/KEY: mat_peptide

(B) LOCATION: 248..1444

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GAATTCGCGG CCGCTGGGAA CGCGGAGCCC CGGACCCGCT CCCGCCGCCT CCGACTCGCC	60
CAGGGGGGGT CGCCGGGAGG AGCCCGGGGG AGAGGGACCA GGAGGGGCCC GCGGCCTCGC	120
AGGGGCGCCC GCGCCCCAC CCCTGCCCCC GCCAGCGGAC CGGTCCCCCA CCCCCGTCC	180
TTCCACC ATG CAC TTG CTG GGC TTC TTC TCT GTG GCG TGT TCT CTG CTC	229
Met His Leu Leu Gly Phe Phe Ser Val Ala Cys Ser Leu Leu	
-20 -15 -10	
GCC GCT GCG CTG CTC CCG GGT CCT CGC GAG GCG CCC GCC GCC GCC GCC	277
Ala Ala Ala Leu Leu Pro Gly Pro Arg Glu Ala Pro Ala Ala Ala Ala	
-5 1 5 10	
GCC TTC GAG TCC GGA CTC GAC CTC TCG GAC GCG GAG CCC GAC GCG GGC	325
Ala Phe Glu Ser Gly Leu Asp Leu Ser Asp Ala Glu Pro Asp Ala Gly	
15 20 25	
GAG GCC ACG GCT TAT GCA AGC AAA GAT CTG GAG GAG CAG TTA CGG TCT	373
Glu Ala Thr Ala Tyr Ala Ser Lys Asp Leu Glu Glu Gln Leu Arg Ser	
30 35 40	
GTG TCC AGT GTA GAT GAA CTC ATG ACT GTA CTC TAC CCA GAA TAT TGG	421
Val Ser Ser Val Asp Glu Leu Met Thr Val Leu Tyr Pro Glu Tyr Trp	
45 50 55	
AAA ATG TAC AAG TGT CAG CTA AGG AAA GGA GGC TGG CAA CAT AAC AGA	469
Lys Met Tyr Lys Cys Gln Leu Arg Lys Gly Gly Trp Gln His Asn Arg	
60 65 70	
GAA CAG GCC AAC CTC AAC TCA AGG ACA GAA GAG ACT ATA AAA TTT GCT	517
Glu Gln Ala Asn Leu Asn Ser Arg Thr Glu Glu Thr Ile Lys Phe Ala	
75 80 85 90	
GCA GCA CAT TAT AAT ACA GAG ATC TTG AAA AGT ATT GAT AAT GAG TGG	565
Ala Ala His Tyr Asn Thr Glu Ile Leu Lys Ser Ile Asp Asn Glu Trp	
95 100 105	
AGA AAG ACT CAA TGC ATG CCA CGG GAG GTG TGT ATA GAT GTG GGG AAG	613
Arg Lys Thr Gln Cys Met Pro Arg Glu Val Cys Ile Asp Val Gly Lys	
110 115 120	

GAG TTT GGA GTC GCG ACA AAC ACC TTC TTT AAA CCT CCA TGT GTG TCC	661
Glu Phe Gly Val Ala Thr Asn Thr Phe Phe Lys Pro Pro Cys Val Ser	
125 130 135	
GTC TAC AGA TGT GGG GGT TGC TGC AAT AGT GAG GGG CTG CAG TGC ATG	709
Val Tyr Arg Cys Gly Gly Cys Cys Asn Ser Glu Gly Leu Gln Cys Met	
140 145 150	
AAC ACC AGC ACG AGC TAC CTC AGC AAG ACG TTA TTT GAA ATT ACA GTG	757
Asn Thr Ser Thr Ser Tyr Leu Ser Lys Thr Leu Phe Glu Ile Thr Val	
155 160 165 170	
CCT CTC TCT CAA GGC CCC AAA CCA GTA ACA ATC AGT TTT GCC AAT CAC	805
Pro Leu Ser Gln Gly Pro Lys Pro Val Ile Ser Phe Ala Asn His	
175 180 185	
ACT TCC TGC CGA TGC ATG TCT AAA CTG GAT GTT TAC AGA CAA GTT CAT	853
Thr Ser Cys Arg Cys Met Ser Lys Leu Asp Val Tyr Arg Gln Val His	
190 195 200	
TCC ATT ATT AGA CGT TCC CTG CCA GCA ACA CTA CCA CAG TGT CAG GCA	901
Ser Ile Ile Arg Arg Ser Leu Pro Ala Thr Leu Pro Gln Cys Gln Ala	
205 210 215	
GCG AAC AAG ACC TGC CCC ACC AAT TAC ATG TGG AAT AAT CAC ATC TGC	949
Ala Asn Lys Thr Cys Pro Thr Asn Tyr Met Trp Asn Asn His Ile Cys	
220 225 230	
AGA TGC CTG GCT CAG GAA GAT TTT ATG TTT TCC TCG GAT GCT GGA GAT	997
Arg Cys Leu Ala Gln Glu Asp Phe Met Phe Ser Ser Asp Ala Gly Asp	
235 240 245 250	
GAC TCA ACA GAT GGA TTC CAT GAC ATC TGT GGA CCA AAC AAG GAG CTG	1045
Asp Ser Thr Asp Gly Phe His Asp Ile Cys Gly Pro Asn Lys Glu Leu	
255 260 265	
GAT GAA GAG ACC TGT CAG TGT GTC TGC AGA GCG GGG CTT CGG CCT GCC	1093
Asp Glu Glu Thr Cys Gln Cys Val Cys Arg Ala Gly Leu Arg Pro Ala	
270 275 280	
AGC TGT GGA CCC CAC AAA GAA CTA GAC AGA AAC TCA TGC CAG TGT GTC	1141
Ser Cys Gly Pro His Lys Glu Leu Asp Arg Asn Ser Cys Gln Cys Val	
285 290 295	
TGT AAA AAC AAA CTC TTC CCC AGC CAA TGT GGG GCC AAC CGA GAA TTT	1189
Cys Lys Asn Lys Leu Phe Pro Ser Gln Cys Gly Ala Asn Arg Glu Phe	
300 305 310	
GAT GAA AAC ACA TGC CAG TGT GTA TGT AAA AGA ACC TGC CCC AGA AAT	1237
Asp Glu Asn Thr Cys Gln Cys Val Cys Lys Arg Thr Cys Pro Arg Asn	
315 320 325 330	
CAA CCC CTA AAT CCT GGA AAA TGT GCC TGT GAA TGT ACA GAA AGT CCA	1285
Gln Pro Leu Asn Pro Gly Lys Cys Ala Cys Glu Cys Thr Glu Ser Pro	
335 340 345	
CAG AAA TGC TTG TTA AAA GGA AAG AAG TTC CAC CAC CAA ACA TGC AGC	1333
Gln Lys Cys Leu Leu Lys Gly Lys Lys Phe His His Gln Thr Cys Ser	
350 355 360	

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TGT TAC AGA CGG CCA TGT ACG AAC CGC CAG AAG GCT TGT GAG CCA GGA      1381
Cys Tyr Arg Arg Pro Cys Thr Asn Arg Gln Lys Ala Cys Glu Pro Gly
      365                      370                      375

TTT TCA TAT AGT GAA GAA GTG TGT CGT TGT GTC CCT TCA TAT TGG AAA      1429
Phe Ser Tyr Ser Glu Glu Val Cys Arg Cys Val Pro Ser Tyr Trp Lys
      380                      385                      390

AGA CCA CAA ATG AGC TAA GATTGTACTG TTTTCCAGTT CATCGATTTT      1477
Arg Pro Gln Met Ser *
395                      400

CTATTATGGA AACTGTGTT GCCACAGTAG AACTGTCTGT GAACAGAGAG ACCCTTGTGG      1537

TCCATGCTAA CAAAGACAAA AGTCTGTCTT TCCTGAACCA TGTGGATAAC TTTACAGGAA      1597

ATGGTACTGG AGCTCATCTG CAAAAGGCCT CTTGTGAAGA CTGGTTTTCT GCCAATGACC      1657

AAACAGCCAA GATTTCCTCT CGTGATTTCT TTAAGAAGAT GACTATATAA TTTATTTCCA      1717

CTAAAAATAT TGTTTCTGCA TTCATTTTTA TAGCAACAAC AATTGGTAAA ACTCACTGTG      1777

ATCAATATTT TTATATCATG CAAAATATGT TTAATAATAA ATGAAAATTG TATTATAAGC      1837

TGCTAAGTTC AGTCCATTAT CATCTTACAT GATGAACGAA AACTACTATC ATGAAGACAC      1897

TGATCTTTCT CTGCCCTTTT TTGTCTCTA ACCAGATGTC ACATATGTAT TACTATGATA      1957

AAAAGTATGA TCCTGTGAAA GAGAGTGTCA GAGGACAACA GAATGCTATT GCTTCATCTC      2017

TTATATGTTT AATGATTATA AACATTTTAG TACATGATAC TTTTGAATTT ATGACCAAGT      2077

GAATCAATAT GAAACATCTT GTAAGATAGA CTAATTAGCA TTGTGATTAA AAGTCATTCA      2137

GTGCTCTGAG AACATTCAGA ATCTTACGTT GGTAGAAAAT CCTGCAGTAT ATATTAAAT      2197

GGCTTTAAAT ATTTTCTCAA AAATAATCTT TTCCAAATAT TTGACTTTTT CTGGCCAGCT      2257

AAAATACTTT TTGTGAGTGG AAGTGCTCCT ATCCAAACAT TTTAAAAAAA GCGCCCGCGA      2317

ATTC                                                                2321

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(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 420 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

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Met His Leu Leu Gly Phe Phe Ser Val Ala Cys Ser Leu Leu Ala Ala
-20          -15          -10          -5

Ala Leu Leu Pro Gly Pro Arg Glu Ala Pro Ala Ala Ala Ala Phe
      1              5              10

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Glu Ser Gly Leu Asp Leu Ser Asp Ala Glu Pro Asp Ala Gly Glu Ala
 15 20 25
 Thr Ala Tyr Ala Ser Lys Asp Leu Glu Glu Gln Leu Arg Ser Val Ser
 30 35 40
 Ser Val Asp Glu Leu Met Thr Val Leu Tyr Pro Glu Tyr Trp Lys Met
 45 50 55 60
 Tyr Lys Cys Gln Leu Arg Lys Gly Gly Trp Gln His Asn Arg Glu Gln
 65 70 75
 Ala Asn Leu Asn Ser Arg Thr Glu Glu Thr Ile Lys Phe Ala Ala Ala
 80 85 90
 His Tyr Asn Thr Glu Ile Leu Lys Ser Ile Asp Asn Glu Trp Arg Lys
 95 100 105
 Thr Gln Cys Met Pro Arg Glu Val Cys Ile Asp Val Gly Lys Glu Phe
 110 115 120
 Gly Val Ala Thr Asn Thr Phe Phe Lys Pro Pro Cys Val Ser Val Tyr
 125 130 135 140
 Arg Cys Gly Gly Cys Cys Asn Ser Glu Gly Leu Gln Cys Met Asn Thr
 145 150 155
 Ser Thr Ser Tyr Leu Ser Lys Thr Leu Phe Glu Ile Thr Val Pro Leu
 160 165 170
 Ser Gln Gly Pro Lys Pro Val Thr Ile Ser Phe Ala Asn His Thr Ser
 175 180 185
 Cys Arg Cys Met Ser Lys Leu Asp Val Tyr Arg Gln Val His Ser Ile
 190 195 200
 Ile Arg Arg Ser Leu Pro Ala Thr Leu Pro Gln Cys Gln Ala Ala Asn
 205 210 215 220
 Lys Thr Cys Pro Thr Asn Tyr Met Trp Asn Asn His Ile Cys Arg Cys
 225 230 235
 Leu Ala Gln Glu Asp Phe Met Phe Ser Ser Asp Ala Gly Asp Asp Ser
 240 245 250
 Thr Asp Gly Phe His Asp Ile Cys Gly Pro Asn Lys Glu Leu Asp Glu
 255 260 265
 Glu Thr Cys Gln Cys Val Cys Arg Ala Gly Leu Arg Pro Ala Ser Cys
 270 275 280
 Gly Pro His Lys Glu Leu Asp Arg Asn Ser Cys Gln Cys Val Cys Lys
 285 290 295 300
 Asn Lys Leu Phe Pro Ser Gln Cys Gly Ala Asn Arg Glu Phe Asp Glu
 305 310 315
 Asn Thr Cys Gln Cys Val Cys Lys Arg Thr Cys Pro Arg Asn Gln Pro
 320 325 330

Leu Asn Pro Gly Lys Cys Ala Cys Glu Cys Thr Glu Ser Pro Gln Lys
 335 340 345
 Cys Leu Leu Lys Gly Lys Lys Phe His His Gln Thr Cys Ser Cys Tyr
 350 355 360
 Arg Arg Pro Cys Thr Asn Arg Gln Lys Ala Cys Glu Pro Gly Phe Ser
 365 370 375 380
 Tyr Ser Glu Glu Val Cys Arg Cys Val Pro Ser Tyr Trp Lys Arg Pro
 385 390 395
 Gln Met Ser *
 400

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- (vii) IMMEDIATE SOURCE:
 - (B) CLONE: FLAG peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Asp Tyr Lys Asp Asp Asp Asp Lys
 1 5

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 191 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (iii) HYPOTHETICAL: NO
- (vii) IMMEDIATE SOURCE:
 - (B) CLONE: Hu VEGF
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Met Asn Phe Leu Leu Ser Trp Val His Trp Ser Leu Ala Leu Leu Leu
 1 5 10 15

Tyr	Leu	His	His	Ala	Lys	Trp	Ser	Gln	Ala	Ala	Pro	Met	Ala	Glu	Gly	
			20				25						30			
Gly	Gly	Gln	Asn	His	His	Glu	Val	Val	Lys	Phe	Met	Asp	Val	Tyr	Gln	
			35				40						45			
Arg	Ser	Tyr	Cys	His	Pro	Ile	Glu	Thr	Leu	Val	Asp	Ile	Phe	Gln	Glu	
			50				55						60			
Tyr	Pro	Asp	Glu	Ile	Glu	Tyr	Ile	Phe	Lys	Pro	Ser	Cys	Val	Pro	Leu	
			65				70						75	80		
Met	Arg	Cys	Gly	Gly	Cys	Cys	Asn	Asp	Glu	Gly	Leu	Gly	Cys	Val	Pro	
			85						90						95	
Thr	Glu	Glu	Ser	Asn	Ile	Thr	Met	Gln	Ile	Met	Arg	Ile	Lys	Pro	His	
			100						105						110	
Gln	Gly	Gln	His	Ile	Gly	Glu	Met	Ser	Phe	Leu	Gln	His	Asn	Lys	Cys	
			115						120						125	
Glu	Cys	Arg	Pro	Lys	Lys	Asp	Arg	Ala	Arg	Gln	Glu	Asn	Pro	Cys	Gly	
			130			135						140				
Pro	Cys	Ser	Glu	Arg	Arg	Lys	His	Leu	Phe	Val	Gln	Asp	Pro	Gln	Thr	
			145			150						155			160	
Cys	Lys	Cys	Ser	Cys	Lys	Asn	Thr	Asp	Ser	Arg	Cys	Lys	Ala	Arg	Gln	
			165						170						175	
Leu	Glu	Leu	Asn	Glu	Arg	Thr	Cys	Arg	Cys	Asp	Lys	Pro	Arg	Arg		
			180						185						190	

(1) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 149 amino acids
(B) TYPE: amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(iii) HYPOTHETICAL: NO

(vii) IMMEDIATE SOURCE:

(B) CLONE: Hu PLGF

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Met	Pro	Val	Met	Arg	Leu	Phe	Pro	Cys	Phe	Leu	Gln	Leu	Leu	Ala	Gly
1				5					10					15	
Leu	Ala	Leu	Pro	Ala	Val	Pro	Pro	Gln	Gln	Trp	Ala	Leu	Ser	Ala	Gly
			20					25					30		
Asn	Gly	Ser	Ser	Glu	Val	Glu	Val	Val	Pro	Phe	Gln	Glu	Val	Trp	Gly
		35					40					45			
Arg	Ser	Tyr	Cys	Arg	Ala	Leu	Glu	Arg	Leu	Val	Asp	Val	Val	Ser	Glu
	50					55				60					

Tyr Pro Ser Glu Val Glu His Met Phe Ser Pro Ser Cys Val Ser Leu
 65 70 75 80
 Leu Arg Cys Thr Gly Cys Cys Gly Asp Glu Asn Leu His Cys Val Pro
 85 90 95
 Val Glu Thr Ala Asn Val Thr Met Gln Leu Leu Lys Ile Arg Ser Gly
 100 105 110
 Asp Arg Pro Ser Tyr Val Glu Leu Thr Phe Ser Gln His Val Arg Cys
 115 120 125
 Glu Cys Arg Pro Leu Arg Glu Lys Met Lys Pro Glu Arg Cys Gly Asp
 130 135 140
 Ala Val Pro Arg Arg
 145

CLAIMS

What is claimed is:

1. An isolated DNA encoding a flk-1 binding protein (flk-1bp), wherein said flk-1bp comprises an amino acid sequence selected from the group consisting of residues -20 to 399 of SEQ ID NO:2 and residues 1 to 399 of SEQ ID NO:2.

2. An isolated DNA of claim 1, wherein said DNA comprises a nucleotide sequence selected from the group consisting of nucleotides 188 to 1444 of SEQ ID NO:1 and nucleotides 248 to 1444 of SEQ ID NO:1.

3. An isolated DNA encoding a human flk-1bp polypeptide, wherein:
a) said flk-1bp polypeptide is characterized by the N-terminal amino acid sequence Gly-Pro-Arg-Glu-Ala-Pro-Ala-Ala-Ala-Ala-Phe-Glu-Ser-Gly- (amino acids 1-15 of SEQ ID NO:2);

b) said flk-1bp polypeptide is capable of binding flk-1; and

c) said flk-1bp polypeptide comprises an amino acid sequence that is at least 90% identical to the amino acid sequence of residues 1 to 399 of SEQ ID NO:2.

4. An isolated DNA encoding an flk-1bp polypeptide capable of binding flk-1, wherein said flk-1bp polypeptide comprises conservative substitution(s) in the amino acid sequence of residues 1 to 399 of SEQ ID NO:2.

5. An isolated DNA encoding a flk-1bp polypeptide, wherein said DNA comprises the flk-1bp cDNA insert of the vector deposited in strain ATCC 69897.

6. An isolated DNA encoding a human flk-1bp polypeptide, wherein a mature form of said flk-1bp polypeptide is characterized by:

a) the N-terminal amino acid sequence Gly-Pro-Arg-Glu-Ala-Pro-Ala-Ala-Ala-Ala-Phe-Glu-Ser-Gly- (amino acids 1-15 of SEQ ID NO:2);

b) the ability to bind flk-1;

c) a calculated molecular weight of 44,823 daltons; and

d) an isoelectric point of 8.678.

7. An expression vector comprising a DNA of any of claims 1 to 6.

8. A process for preparing a flk-1bp polypeptide, comprising culturing a host cell transformed with a vector of claim 7 under conditions promoting expression of flk-1bp, and recovering the flk-1bp polypeptide.

9. A purified flk-1bp polypeptide, wherein:

a) said flk-1bp polypeptide is characterized by the N-terminal amino acid sequence Gly-Pro-Arg-Glu-Ala-Pro-Ala-Ala-Ala-Ala-Phe-Glu-Ser-Gly- (amino acids 1-15 of SEQ ID NO:2);

b) said flk-1bp polypeptide is capable of binding flk-1; and

c) said flk-1bp polypeptide comprises an amino acid sequence that is at least 90% identical to the amino acid sequence of residues 1 to 399 of SEQ ID NO:2.

10. A purified flk-1bp polypeptide of claim 9, comprising the amino acid sequence presented as residues 1 to 399 of SEQ ID NO:2.

11. A purified human flk-1bp protein encoded by the cDNA insert of the recombinant vector deposited in strain ATCC 69897.

12. A purified mature human flk-1bp protein characterized by:

a) the N-terminal amino acid sequence Gly-Pro-Arg-Glu-Ala-Pro-Ala-Ala-Ala-Ala-Phe-Glu-Ser-Gly- (amino acids 1-15 of SEQ ID NO:2);

b) the ability to bind flk-1;

c) a calculated molecular weight of 44,823 daltons; and

d) an isoelectric point of 8.678.

13. A purified flk-1bp polypeptide, wherein said flk-1bp polypeptide comprises conservative substitution(s) in the amino acid sequence of residues 1 to 399 of SEQ ID NO:2, and is capable of binding flk-1.

14. An oligomer comprising from two to four flk-1bp polypeptides according to claim 9.

15. An oligomer of claim 14, comprising two flk-1bp polypeptides.

FIGURE 1

Hu VEGF	lllylhakwsqaapmaegggqnhhevkvfmdv-----
Hu PLGF	aglalpavppqqwalsagnssevvpfgev-----
Hu flk-1bp	aaallpgpreapaaaaafesgldlsdaepdageatatayaskdleeql
Hu VEGF	-----
Hu PLGF	-----
Hu flk-1bp	rsvssvdelmtvlypeywkmykcqlrkqgqwhnreqanlnsrteetikfaaaahynteilk
Hu VEGF	-----YQRSY CHPI ETLV DIFQ EYPDEIEYI FKPSQV PLM RC GGGC ND EG EG VPTEESN
Hu PLGF	-----WGRSY CRAL RRLV DVVS EYPSEVEHM CSPSQV SLL RC TGCC GD EN LH QVPVETAN
Hu flk-1bp	sidnewrktq CMPR EVC I DVGK EFGVATNTF EKPPQV SVX RC GGGC NS EG EG CMNTSTSY
Hu VEGF	ITMQIMR IKPH--QGQHIGEM GFLQHNK CECRPKKDRARQenpcgp-----
Hu PLGF	VTMQLLK I RSGDRPSYVELTF SQHVRC CRplrekmkpercgdavn Er-----
Hu flk-1bp	LSKTLFE ITVPLSQGPKPVTI: SFANHTS CRMSKLDVYRQvhsilr Fslpatlpqcqaan
Hu VEGF	-----
Hu PLGF	-----
Hu flk-1bp	ktcptnymwnnhicrcJagedmfssdagddstdgfhdicgpnkeldeetcgcvcraglr
Hu VEGF	-----
Hu PLGF	-----
Hu flk-1bp	pascgphkeldrnscqcvcknklfpsqcganrefdentcgcq CERRKHLFVQD PQTC KC
Hu VEGF	-----
Hu PLGF	-----
Hu flk-1bp	SCKNTDSRCARQLELNER TCR CDKPRR-----
Hu VEGF	-----
Hu PLGF	-----
Hu flk-1bp	ECTESPQKCLLKGGKFFHQ TCS yrrpctnrqacepgfsyseevcvcvpsywkripqms

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/17584**A. CLASSIFICATION OF SUBJECT MATTER**

IPC(6) : C12N 15/12, 15/67, 15/70, 15/79; C07K 14/00, 14/47

US CL : 536/23.5; 435/320.1, 69.1; 530/350, 402

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 536/23.5; 435/320.1, 69.1; 530/350, 402

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

APS/USPAT; STN/MEDLINE, HCAPLUS

search terms: VEGF, vascular endothelial growth factor, flk 1

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	WO 95/24473 A1 (HUMAN GENOME SCIENCES, INC.) 14 September 1995 (14.09.95), see entire document, especially Figure 1.	1-15
A	EP 0 506 477 A1 (MERCK & CO., INC.) 30 September 1992 (30.09.92), see entire document, especially Figure 1.	1-15
P, X	LEE et al. Vascular endothelial growth factor-related protein: A ligand and specific activator of the tyrosine kinase receptor Flt4. Proceedings of the National Academy of Science USA, March 1996, Vol. 93, pages 1988-1992, especially page 1990, Figure 1.	1-13

☒ Further documents are listed in the continuation of Box C.
 ☐ See patent family annex.

* Special categories of cited documents:	*T later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
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*L document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*Z document member of the same patent family
*O document referring to an oral disclosure, use, exhibition or other means	
*P document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search

12 FEBRUARY 1997

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INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/17584

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P, X	JOUKOV et al. A Novel Vascular Endothelial Growth Factor, VEGF-C, is a Ligand for the Flt4 (VEGFR-3) and KDR (VEGFR-2) Receptor Tyrosine Kinases. EMBO Journal, Volume 15, Number 2, pages 290-298, especially page 293, Figure 3.	1-13